

F16.2

[illegible]

420
430
440

---

SEQ ID NO: 5     A C T G A C C C C T T T T

↓   ↓                ↓   ↓                ↓   ↓  
 G   A                T   T                A   A

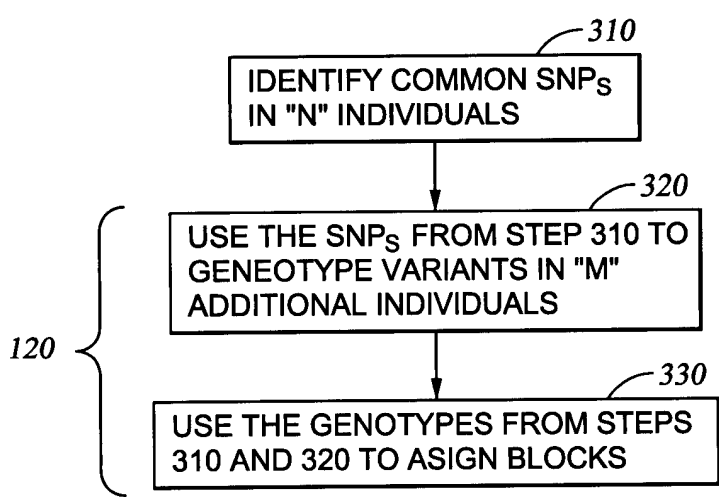
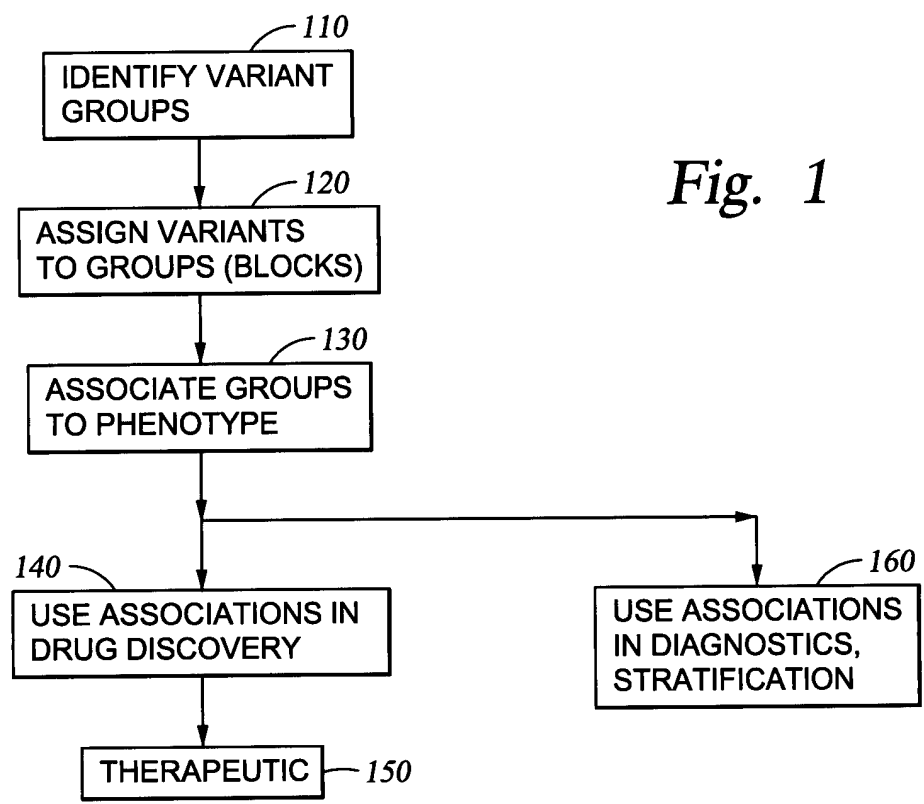
SEQ ID NO: 6	... A <sup>✓</sup> G T <sup>✓</sup> A A C <sup>✓</sup> C C <sup>✓</sup> C T T <sup>✓</sup> T T <sup>✓</sup> ...	} 410
SEQ ID NO: 7	... A C T G A C C C C T T T T ...	
SEQ ID NO: 8	... A C T G A C T C T T T A A ...	

}
420

G ... A ... T ... T ... T T

F16. 4



BLOCK: A CHROMOSOMAL REGION  
WHICH TENDS TO BE INHERITED AS A UNIT,  
WITH A SMALL NUMBER OF COMMON FORMS

	241	242	243	244	245	246	247	248	249	250	251	252	253
SEQ ID NO:1	W {	...A...	G...	A...	T...	T...	C...	G...	A...	T...	A...	A...	C...
SEQ ID NO:2	X {	...A...	G...	A...	C...	T...	A...	C...	A...	T...	A...	A...	C...
SEQ ID NO:3	Y {	...T...	A...	T...	T...	T...	C...	G...	A...	T...	A...	A...	C...
SEQ ID NO:4	Z {	...T...	A...	T...	C...	T...	A...	C...	A...	A...	T...	C...	A...
		261			262					263			

Fig. 2

Fig. 4